RiboV/es² June 10th-13th 2018

Where RNA Minds Reunite



Lethbridge

Waterton



SPEAKERS

Martin Hirst, University of British Columbia, Canada Steven Jones, Canada's Michael Smith Genome Sciences Centre, Canada Jennifer Kugel, University of Colorado Boulder, USA Michelle Scott, Université de Sherbrooke, Canada Nahum Sonenberg, McGill University, Canada

SPECIAL FOCUS

RNA in Health and Disease New RNA Technologies

The organizers thank the following sponsors for their kind support of the 2018 RiboWest Conference































RiboWest 2018 Program

Sunday, June 10, 2018

7:30 am – 5:00 pm	Trip to Waterton Lakes National Park with networking lunch	
5:00 – 7:00 pm	Registration with light snacks	Markin Hall Atrium
5:45 – 6:00 pm	Opening Remarks: Dr. Erasmus Okine (Vice President Research, University of Lethbridge)	Markin Hall Atrium
6:00 – 7:00 pm	Gairdner Lecture: Dr. Nahum Sonenberg	Markin Hall Atrium
7:00 – 8:00 pm	Alberta Epigenetics Network (AEN) sponsored Opening Keynote Lecture: Dr. Steven Jones	Markin Hall Atrium
8:00 – 10:00 pm	Welcome Reception and Networking Opportunity	Lux Hotel Lobby
Monday, June 11, 2018		
7:00 – 8:00 am	Breakfast	Markin Hall Atrium
8:00 – 8:45 am	AEN Sponsored "Breakthrough in Epigenetics" Lecture: Dr. Martin Hirst	Markin Hall Atrium
9:00 – 10:20 am	Session I: Translation Regulation	PE261
10:20 – 10:45 am	Coffee Break	Markin Hall Atrium
11:00 am – 12:15 pm	Session II: RNA in Infectious Diseases	PE261
12:15 – 1:15 pm	Lunch	Markin Hall Atrium
1:15 – 1:35 pm	Group Photo	Outside of Markin Hall
1:45 – 3:20 pm	Session III: Gene Expression Regulation and Diseases	PE261
3:20 – 3:40 pm	Coffee Break	Markin Hall Atrium

3:40 – 4:30 pm	Panel Discussion on Grant Writing and Team Building for Tri-Council Grants	Markin Hall Atrium
4:30 – 5:00 pm	Beckman Seminar	Markin Hall Atrium
5:00 – 6:30 pm	Poster Session Part 1 (even-numbered posters)	Markin Hall
6:30 – 8:00 pm	Dinner and Networking session	Markin Hall Atrium
8:30 – 10:30 pm	Informal social	Off campus
Tuesday, June 12, 2018		
7:00 – 8:00 am	Breakfast	Markin Hall Atrium
8:00 – 8:45 am	Invited RiboClub Lecture: Dr. Michelle Scott	Markin Hall Atrium
9:00 – 10:35	Session IV: RNA Structure-Function	PE261
10:35 – 11:00 am	Coffee Break	Markin Hall Atrium
11:15 am – 12:45 pm	Session V: Computational Biology and	PE261
11.13 diii 12.43 piii	Transcriptomics	1 1 2 0 1
12:45 – 1:45 pm		Markin Hall Atrium
·	Transcriptomics	Markin Hall Atrium
12:45 – 1:45 pm	Transcriptomics Lunch Concurrently: Principal investigators discussion	Markin Hall Atrium
12:45 – 1:45 pm 1:25 – 1:45 pm	Transcriptomics Lunch Concurrently: Principal investigators discussion on the venue for RiboWest Conference 2019	Markin Hall Atrium Markin Hall Atrium
12:45 – 1:45 pm 1:25 – 1:45 pm 2:00 – 2:45 pm	Transcriptomics Lunch Concurrently: Principal investigators discussion on the venue for RiboWest Conference 2019 Session VI: Synthetic Biology and Biophysics	Markin Hall Atrium Markin Hall Atrium PE261
12:45 – 1:45 pm 1:25 – 1:45 pm 2:00 – 2:45 pm 2:45 – 3:10 pm	Transcriptomics Lunch Concurrently: Principal investigators discussion on the venue for RiboWest Conference 2019 Session VI: Synthetic Biology and Biophysics Coffee Break Poster Session Part 2	Markin Hall Atrium Markin Hall Atrium PE261 Markin Hall Atrium
12:45 – 1:45 pm 1:25 – 1:45 pm 2:00 – 2:45 pm 2:45 – 3:10 pm 3:15 – 4:45 pm	Transcriptomics Lunch Concurrently: Principal investigators discussion on the venue for RiboWest Conference 2019 Session VI: Synthetic Biology and Biophysics Coffee Break Poster Session Part 2 (odd-numbered posters)	Markin Hall Atrium Markin Hall Atrium PE261 Markin Hall Atrium Markin Hall

RiboWest 2018

Wednesday, June 13, 2018

7:30 – 8:30 am	Breakfast	Markin Hall Atrium
8:30 – 10:30 am	Session VII: AEN Sponsored Young Investigator Session	PE261
10:30 – 11:00 am	Conference wrap up, Coffee, and Departure	Markin Hall Atrium

Oral Presentations

Sunday, June 10, 2	2018	
	Nahum Sonenberg (McGill University, QC, Canada) Okine and Dr. Nehal Thakor	
6:00 – 7:00 pm	Diverse Functions of the Cap-binding Protein 4EHP-Complex in Control of mRNA Translation	1
Sciences Centre, BC C	ning Keynote Lecture: Dr. Steven Jones (Canada's Michael Smith Gen Cancer Agency, BC, Canada) Okine and Dr. Athan Zovoilis	ome
7:00 – 8:00 pm	Using genomics and transcriptomics to develop a platform for precision medicine in oncology	2
Monday, June 11, Focus: RNA in Hea		
AEN Sponsored "Brea Columbia, BC, Canad Chair: Dr. Tommy A	•	tish
8:00 – 8:45 am	Exploiting cis-regulatory enhancers as therapeutic targets in cancer	3
Session I: Translation Chair: Dr. Nehal Tha		
9:00 – 9:20 am	Poul Sorensen RNA stress granules mediate translational control of tumor metastasis	37
9:20 – 9:35 am	Joseph Ross Eukaryotic initiation factor 5B (eIF5B) provides a critical cell survival switch to glioblastoma cells via regulation of apoptosis	32
9:35 – 9:50 am	Tyson Graber Virus-induced 5' UTR switching drives translation of an mRNA	17

infection efficacy

encoding an inositol phosphatase that limits oncolytic virus

9:50 – 10:05 am	Seyed Mehdi Jafarnejad Regulation of ERK signalling pathway by a microRNA and mRNA translation machinery	20
10:05 – 10:20 am	Kamiko Bressler The role of eukaryotic initiation factor 5B (eIF5B) in upstream open reading frame (uORF)-mediated translation	11
Session II: RNA in Infect	ious Diseases	
Chair: Dr. Chris Nelson		
11:00 – 11:15 am	Mamata Panigrahi Altered Translation Regulation Promotes miR-122- independent Replication of Hepatitis C Virus Genomes	29
11:15 – 11:30 am	Monica Mesa Perez Dissecting the functions of a cytoplasmic isoform of ARS2 in RNA metabolism	27
11:30 – 11:45 am	Tyler Mrozowich Towards Obtaining a Nanoscale Structure of the 3' Terminal Region of Japanese Encephalitis Virus Genome	28
11:45 am – 12:00 pm	Jibin Sadasivan Cricket Paralysis Virus 1A protein disrupts stress granule formation	33
12:00 – 12:15 pm	Rasika Kunden Mapping the region on the Hepatitis C virus genome to which miR-122 and other small RNA annealing promotes virus replication	24
Session III: Gene Expres	sion Regulation and Diseases	
Chair: Dr. Richard Fahl	man	
1:45 – 2:05 pm	Hasan Uludag siRNA Therapy of Chronic Myeloid Leukemia	38
2:05 – 2:20 pm	Logan Brand I get knocked down, but I get up again: transcription and 5' UTR contribution to temperature-dependent regulation of a cyanobacterial RNA helicase	10

2:20 – 2:35 pm	Neda Savic Paralogous <i>S. cerevisiae</i> histone chaperones Fpr3 and Fpr4 regulate transcription at rDNA and at select protein coding genes	34
2:35 – 2:50 pm	Sean Ritter Protein Abundance of the RNA Helicase CrhR is Regulated via Convergent Sensing of the Redox Poise of the Electron Transport Chain	30
2:50 – 3:05 pm	Dani Kalsbeek DNA damage checkpoint adaptation leads to chromosome loss and rearrangement in cancer cells	22
3:05 – 3:20 pm	John Maringa Githaka Non-canonical role of the Bcl-2 protein BAD in translation and mammary gland morphogenesis	16
Beckman Lecture: Bo Chair: Dr. Trushar P	ob Rodriguez (Beckman Coulter Inc.) Patel	
4:30 – 5:00 pm	Analytical Ultracentrifugation: Overview and Common Applications	6
Tuesday, June 12,	, 2018	
Focus: New RNA	Technologies	
Invited RiboClub Lec Chair: Dr. Stacey W	ture: Dr. Michelle Scott (Université de Sherbrooke, QC, Canada) etmore	
8:00 – 8:45 am	A pan-transcriptome view of mid-size RNAs	4
Session IV: RNA Stru Chair: Dr. Ute Koth		
9:00 – 9:20 am	Savraj Grewel tRNA synthesis and the regulation of growth in <i>Drosophila</i>	18
9:20 – 9:35 am	Emily Soon TrmA is a tRNA modification enzyme that functions as a tRNA chaperone <i>in vitro</i> and <i>in vivo</i>	36

9:35 – 9:50 am	Sarah Schultz Hierarchical modification of tRNA by TrmA, TruB, and TrmB	35
9:50 – 10:05 am	Mathew Kirby A short linker region of factorless internal ribosome entry site mediates ribosomal positioning and translational activity	23
10:05 – 10:20 am	Luc Roberts Evidence against RNA structure-based translation initiation in bacteria	31
10:20 – 10:35 am	Beatrice Fung Elucidation of 3' exon requirements for a novel group II intron	14
Session V: Computation	nal Biology and Transcriptomics	
Chair: Dr. Sean McKen		
11:15 – 11:30 am	Chris Isaac Abnormal SINE B2 RNA processing is associated with amyloid and brain aging pathology	19
11:30 – 11:45 am	Gabrielle Deschamps-Francoeur CoCo: RNA-seq Read Assignment Correction for Nested Genes and Multimapped Reads	12
11:45 am – 12:00 pm	Dylan Girodat Conserved Dynamics of the D2 Dopamine Receptor Transmembrane Helix 3 Reveal an Activation Pathway Amongst G-Protein Coupled Receptors	15
12:00 – 12:15 pm	Katie Wilson A Survey of RNA-Protein π -Interactions: A Comparison of Natural Occurrences and Structures, and Computationally Predicted Strengths	40
12:15 – 12:30 pm	Mohamed Aboelnga Computational insights into the multifunctional AP endonuclease: Establishing a novel phosphodiester bond cleavage mechanism	7
12:30 – 12:45 pm	Olivia Marasco Testing the Vattem-Wek Hypothesis for ATF4 translation regulation by upstream open reading frames using <i>in silico</i> and <i>in cellulo</i> methods	25

Session VI: Synthetic Biology and Biophysics		
Chair: Dr. Greg Mori	n	
2:00 – 2:15 pm	Justin Vigar Non-Canonical Translation Initiation Devices for Engineering Biology	39
2:15 – 2:30 pm	Ewan McRae The C-terminus of DDX21 recognizes the 2'OH of loop nucleotides in RNA guanine quadruplexes	26
2:30 – 2:45 pm	Simmone D'souza Interaction Studies of DDX17 with Rift Valley Fever Virus and Japanese Encephalitis Virus Non-coding RNA	13
Closing Keynote Lectu Chair: Dr. Athan Zov	ure: Dr. Jennifer Kugel (University of Colorado Boulder, CO, USA) oilis	
4:45 – 5:45 pm	Regulation of RNA polymerase II by SINE encoded ncRNAs	5
Session VII: Young Inv Chair: Dr. Raja Singh		
8:30 – 9:00 am	Athanasios Zovoilis Epigenetic deregulation through abnormal processing of SINE non-coding RNAs in aging brain and dementia: An integrative RNA genomics approach	41
9:00 – 9:30 am	Tim Audus Uncovering the Mechanisms Regulating Physiological Amyloid Aggregation	8
9:30 – 10:00 am	Andriy Bilichak Genome editing in wheat microspores and haploid embryos mediated by delivery of ZFN proteins and cell penetrating peptide complexes	9
10:00 – 10:30 am	Darryl Jones Automated Phylogenomics For Carbohydrate Active Enzyme Discovery From Animal Microbiomes	21